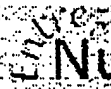


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Links

wk22d07.x1 NCI_CGAP_Lym12 Homo sapiens cDNA clone IMAGE:2413069 3' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|5436445|gb|AI817366.1|5436445]

☐ 2: [AI797688](#) Reports

Links

we90d08.x1 Soares_NFL_T_GBC_S1 Homo sapiens cDNA clone IMAGE:2348367 3' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|5363160|gb|AI797688.1|5363160]

☐ 3: [AI794615](#) Reports

Links

fc46f02.y1 Zebrafish WashU MPIMG EST Danio rerio cDNA clone IMAGE:3724443 5' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|5342331|gb|AI794615.1|5342331]

☐ 4: [AI762942](#) Reports

Links

wh94f09.x1 NCI_CGAP_CLL1 Homo sapiens cDNA clone IMAGE:2388425 3' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|5178609|gb|AI762942.1|5178609]

☐ 5: [AI726943](#) Reports

Links

BNLGHi6897 Six-day Cotton fiber Gossypium hirsutum cDNA 5' similar to (AB012122) TIP49 [Homo sapiens] gi|3243035, MRNA sequence
gi|5045795|gb|AI726943.1|5045795]

☐ 6: [AI725799](#) Reports

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BNLGHi13195 Six-day Cotton fiber Gossypium hirsutum cDNA 5' similar to (AB012122) TIP49 [Homo sapiens] gi|3243035, MRNA sequence
gi|5044651|gb|AI725799.1|5044651]

☐ 7: [AI722028](#) Reports

Links

fd18e04.y1 Sugano Kawakami zebrafish DRA Danio rerio cDNA clone IMAGE:2351934 5' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|5040357|gb|AI722028.1|5040357]

☐ 8: [AI721333](#) Reports

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fd18e04.x1 Sugano Kawakami zebrafish DRA Danio rerio cDNA clone IMAGE:2351934
3' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|5039662|gb|AI721333.1|[5039662]

☐ 9: AB002406 Reports

Links

Rattus norvegicus mRNA for TIP49, complete cds
gi|2225876|dbj|AB002406.1|[2225876]

☐ 10: AI672922 Reports

Links

we74b05.x1 Soares_Dieckgraefe_colon_NHCD Homo sapiens cDNA clone
IMAGE:2346801 3' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|4852653|gb|AI672922.1|[4852653]

☐ 11: AI659383 Reports

Links

tul1f02.x1 NCI_CGAP_Pr28 Homo sapiens cDNA clone IMAGE:2250747 3' similar to
TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|4762953|gb|AI659383.1|[4762953]

☐ 12: AB014075 Reports

Links

Clostridium histolyticum genes for hypoxanthine-guanine phosphoribosyl-transferase
(HGPRTase), GTPase and 12 ORFs, complete and partial cds
gi|3868863|dbj|AB014075.1|[3868863]

☐ 13: B88467 Reports

Links

CpG0213B CpIOWAgDNA1 Cryptosporidium parvum genomic similar to TBP(TATA-
binding protein)-interacting protein 49 (TIP49) homolog, genomic survey sequence
gi|2970893|gb|B88467.1|[2970893]

☐ 14: AI591696 Reports

Links

vq30h01.y1 Barstead stromal cell line MPLRB8 Mus musculus cDNA clone
IMAGE:1095793 5' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|4600744|gb|AI591696.1|[4600744]

☐ 15: D84298 Reports

Links

Deinococcus radiodurans DNA for ruvB protein, complete cds
gi|4586353|dbj|D84298.1|[4586353]

☐ 16: AE000613 Reports

Links

Helicobacter pylori 26695 section 91 of 134 of the complete genome
gi|2314200|gb|AE000613.1|AE000613[2314200]

☐ 17: AI585221 Reports

Links

fb95g04.y1 Zebrafish WashU MPIMG EST Danio rerio cDNA clone IMAGE:3719670 5'
similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|4571118|gb|AI585221.1|[4571118]

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- ☐ 18: [AI584574](#) Reports Links
fb95g04.x1 Zebrafish WashU MPIMG EST Danio rerio cDNA clone IMAGE:3719670 3' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|4570471|gb|AI584574.1|[4570471]
- ☐ 19: [AI565821](#) Reports Links
tn20g08.x1 NCI_CGAP_Brn25 Homo sapiens cDNA clone IMAGE:2168222 3' similar to SW:YEBA_ECOLI P24204 HYPOTHETICAL 46.7 KD PROTEIN IN MSBB-RUVB INTERGENIC REGION ;, MRNA sequence
gi|4524278|gb|AI565821.1|[4524278]
- ☐ 20: [AI539151](#) Reports Links
tp73c06.x1 NCI_CGAP_Ut3 Homo sapiens cDNA clone IMAGE:2204938 3' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|4453286|gb|AI539151.1|[4453286]
- ☐ 21: [AI503510](#) Reports Links
vq30h01.x1 Barstead stromal cell line MPLRB8 Mus musculus cDNA clone IMAGE:1095793 3' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|4401361|gb|AI503510.1|[4401361]
- ☐ 22: [AI421332](#) Reports Links
tf14f12.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:2096207 3' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|4267263|gb|AI421332.1|[4267263]
- ☐ 23: [AI298410](#) Reports Links
qn01f05.x1 NCI_CGAP_Lu5 Homo sapiens cDNA clone IMAGE:1896993 3' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|3958146|gb|AI298410.1|[3958146]
- ☐ 24: [AF064953](#) Reports Links
Coxiella burnetii Holliday junction DNA helicase RuvB gene, complete cds
gi|3248954|gb|AF064953.1|AF064953[3248954]
- ☐ 25: [AE001471](#) Reports Links
Helicobacter pylori, strain J99 section 32 of 132 of the complete genome
gi|4154880|gb|AE001471.1|AE001471[4154880]
- ☐ 26: [AI361999](#) Reports Links
qy49c08.x1 NCI_CGAP_Brn23 Homo sapiens cDNA clone IMAGE:2015342 3' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|4113620|gb|AI361999.1|[4113620]
- ☐ 27: [AI351057](#) Reports Links
qt22c06.x1 NCI_CGAP_GC4 Homo sapiens cDNA clone IMAGE:1948330 3' similar to

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gi|4088263|gb|AI351057.1|[4088263]

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- ☐ 28: [AF070735](#) Reports
Homo sapiens RuvB-like protein RUVBL1 mRNA, complete cds
gi|3243034|gb|AF070735.1|AF070735[3243034]

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- ☐ 29: [AI084169](#) Reports
qt27e09.x1 NCI_CGAP_Bm25 Homo sapiens cDNA clone IMAGE:1751272 3' similar to
TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3422592|gb|AI084169.1|[3422592]

Links

- ☐ 30: [AI183800](#) Reports
qc18h08.x1 Soares_fetal_lung_NbHL19W Homo sapiens cDNA clone IMAGE:1739391 3'
similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3734438|gb|AI183800.1|[3734438]

Links

- ☐ 31: [AI096984](#) Reports
qb42d09.x1 NCI_CGAP_Bm23 Homo sapiens cDNA clone IMAGE:1702769 3' similar to
TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3446566|gb|AI096984.1|[3446566]

Links

- ☐ 32: [AI078327](#) Reports
oz12d03.x1 Soares_fetal_liver_spleen_1NFLS_S1 Homo sapiens cDNA clone
IMAGE:1675109 3' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3412735|gb|AI078327.1|[3412735]

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- ☐ 33: [U55214](#) Reports
Treponema pallidum RuvB' gene, partial cds, and TroA, TroB, TroC, TroD, TroR, Pgm,
Pfs, and Tpp15 genes, complete cds
gi|1777931|gb|U55214.1|TPU55214[1777931]

Links

- ☐ 34: [AI033780](#) Reports
ox20a01.x1 Soares_fetal_liver_spleen_1NFLS_S1 Homo sapiens cDNA clone
IMAGE:1656840 3' similar to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3254733|gb|AI033780.1|[3254733]

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- ☐ 35: [AA906726](#) Reports
ok78g03.s1 NCI_CGAP_GC4 Homo sapiens cDNA clone IMAGE:1520116 3' similar to
TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3042312|gb|AA906726.1|[3042312]

Links

- ☐ 36: [AI049841](#) Reports
an21e02.x1 Gessler Wilms tumor Homo sapiens cDNA clone IMAGE:1699322 3' similar
to TR:O35753 O35753 TIP49. ;, mRNA sequence
gi|3298958|gb|AI049841.1|[3298958]

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- ☐ 37: [AA947109](#) Reports Links
om79a04.s1 NCI_CGAP_Kid3 Homo sapiens cDNA clone IMAGE:1553358 3' similar to
TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|3110504|gb|AA947109.1|[3110504]
- ☐ 38: [AA974522](#) Reports Links
of13d01.s1 NCI_CGAP_Co12 Homo sapiens cDNA clone IMAGE:1420993 3' similar to
TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|3149702|gb|AA974522.1|[3149702]
- ☐ 39: [AI006350](#) Reports Links
ua70f09.r1 Soares_thymus_2NbMT Mus musculus cDNA clone IMAGE:1362857 5'
similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|3215959|gb|AI006350.1|[3215959]
- ☐ 40: [AA985767](#) Reports Links
ua66e06.r1 Soares_thymus_2NbMT Mus musculus cDNA clone IMAGE:1362466 5'
similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|3167156|gb|AA985767.1|[3167156]
- ☐ 41: [AB012122](#) Reports Links
Homo sapiens mRNA for TIP49, complete cds
gi|3132307|dbj|AB012122.1|[3132307]
- ☐ 42: [AA878287](#) Reports Links
oc57g05.s1 NCI_CGAP_Lu5 Homo sapiens cDNA clone IMAGE:1415768 3' similar to
TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|2987252|gb|AA878287.1|[2987252]
- ☐ 43: [AA869108](#) Reports Links
vq30h01.r1 Barstead stromal cell line MPLRB8 Mus musculus cDNA clone
IMAGE:1095793 5' similar to TR:O35753 O35753 TIP49. ;, MRNA sequence
gi|2964553|gb|AA869108.1|[2964553]
- ☐ 44: [AA606258](#) Reports Links
vo47c01.r1 Barstead mouse irradiated colon MPLRB7 Mus musculus cDNA clone
IMAGE:1053024 5' similar to SW:RUVB_HAEIN P44631 HOLLIDAY JUNCTION DNA
HELICASE RUVB. ;, MRNA sequence
gi|2454716|gb|AA606258.1|[2454716]
- ☐ 45: [D90829](#) Reports Links
E.coli genomic DNA, Kohara clone #337(41.9-42.3 min.)
gi|1736504|dbj|D90829.1|[1736504]
- ☐ 46: [D90828](#) Reports Links
E.coli genomic DNA, Kohara clone #336gap(41.6-41.9 min.)
gi|1736490|dbj|D90828.1|[1736490]

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- ☐ 47: D83138 Reports [Links](#)
Pseudomonas aeruginosa *ruvC*, *ruvA* and *ruvB* genes for Holliday junction-related proteins
gi|1183837|dbj|D83138.1|PSEUUVABC[1183837]
- ☐ 48: U52110 Reports [Links](#)
Sulfolobus solfataricus putative ribokinase and *Dbh* genes, complete cds, and putative
ATPase gene, partial cds
gi|1706950|gb|U52110.1|SSU52110[1706950]
- ☐ 49: U38840 Reports [Links](#)
Thermotoga maritima *RuvB* protein (RUVB) gene, complete cds
gi|1063667|gb|U38840.1|TMU38840[1063667]
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E. coli *ruv* region
gi|42901|emb|X07091.1|ECRUUVAB[42901]
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Synechocystis sp. *ruvB* gene, complete cds, and *secA* gene, partial cds
gi|1256587|gb|U38892.1|SSU38892[1256587]
- ☐ 52: M86239 Reports [Links](#)
Cyanophora paradoxa cyanelle transfer RNA-Asn (*trnN*) gene, 3' end; transfer RNA-Leu
(*trnL*) gene; FA/FB (*psaC*) gene, complete cds; ORF243, complete cds; ORF162, 3' end
gi|336634|gb|M86239.1|CPACYLNAA[336634]
- ☐ 53: M21298 Reports [Links](#)
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sensitivity to UV radiation, complete cds
gi|147781|gb|M21298.1|ECORUVABA[147781]
- ☐ 54: U00011 Reports [Links](#)
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